

STIC Search Report Biotech-Chem Library

STIC Database Tracking Number: 103763

TO: Terra Gibbs

Location: CM1/12A12&11E12

Art Unit: 1635

Search Notes

Tuesday, September 23, 2003

Case Serial Number: 10/008789

From: David Schreiber

Location: Biotech-Chem Library

CM1-6A03

Phone: 308-4292

david.schreiber@uspto.gov

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From: Sent:

Gibbs, Terra Thursday, September 11, 2003 5:38 PM STIC-Biotech/ChemLib

To: Subject: Sequence search request...

F. TOENVED

SEP 12 2033

. LCH/CREM. DIVISION (STIC)

Could you please search USSN 10/008,789 SEQ ID NO:3 for the following sequence:

5'-GACTCAGCAGTCAGTGGTGACGGTGGC-3'

Also can you check SEQ ID NO:3 for the complement of the above sequence?

Terra Cotta Gibbs, Ph.D. Art Unit 1635 CM1, 12A12 703-306-3221

Searcher:	1. Sohve. Der
Phone:	308-4292
Location:	CONT GAOS
Date Picked U	p:
Date Complete	ed: 9123
Searcher Prep	/Review:/_3_
Clerical:	
Online time:	5.(

TYPE OF SEARCH:
NA Sequences: >
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

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VENDOR/COST (where applic.)	
STN:	
DIALOG:	
Questel/Orbit:	
DRLink:	
Lexis/Nexis:	
Sequence Sys.: Co., as c	ł.
WWW/Internet:	_
Other (specify):	



STIC SEARCH RESULTS FEEDBACK FORM

Biotech-Chem Library

Questions about the scope or the results of the search? Contact the searcher or contact:

Mary Hale, Information Branch Supervisor 308-4258, CM1-1E01

Voluntary Results Feedback Form
> I am an examiner in Workgroup: Example: 1610
> Relevant prior art found, search results used as follows:
102 rejection
☐ 103 rejection
☐ Cited as being of interest.
Helped examiner better understand the invention.
Helped examiner better understand the state of the art in their technology.
Types of relevant prior art found:
☐ Foreign Patent(s)
Non-Patent Literature (journal articles, conference proceedings, new product announcements etc.)
> Relevant prior art not found:
Results verified the lack of relevant prior art (helped determine patentability).
Results were not useful in determining patentability or understanding the invention.
Comments:

Drop of for send completed toms to STC/Blotech-Chem Library GM1 ≒Che. Desk



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GenCore version 5.1.6 Copyright (c) 1993 - 2003 Compugen Ltd.
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September 23, 2003, 07:46:57; Search time 0.001 Seconds (without alignments) 94.770 Million cell updates/sec
OM nucleic - nucleic search, using sw model
                                                        Run on:
```

FRAGMENT

1 gactcagcagtcagtggtgacggtggc 27 Title: Perfect score: Sequence:

IDENTITY_NUC Gapext 0.5 Scoring table:

~ Total number of hits satisfying chosen parameters:

1 seqs, 1755 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

esult Query No. Score Match Length DB ID Description	
ID	27 100.0 1755 1 us-10-008-789-3 0.6 39.3 1755 1 us-10-008-789-3
98	
Length	1755 1755
Query	27 100.0 10.6 39.3
Score Match Length DB ID	10.6
o t	77
Result No.	υ

ALIGNMENTS

Gaps ö Query Match 100.0%; Score 27; DB 1; Length 1755; Best Local Similarity 100.0%; Pred. No. 0; Matches 27; Conservative 0; Mismatches 0; Indels RESULT 1 us-10-008-789-3/c

ö

11993 GACTCAGCAGTCAGTGACGTGGC 1567 1 GACTCAGCAGTCAGTGGTGACGGTGGC 27 ò

g

RESULT 2 us-10-008-789-3

Gaps ö Query Match 39.3%; Score 10.6; DB 1; Length 1755; Best Local Similarity 64.0%; Pred. No. 0; Matches 16; Conservative 0; Mismatches 9; Indels 0;

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402 CCTGCAGACAGGGGGGCCTTCGC 426 3 CTCAGCAGTCAGTGGTGACGGTGGC 27 ò 셤

Search completed: September 23, 2003, 07:46:57 Job time : 0.001 secs

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 2 summaries

us-10-008-789-3:*